



#15/A

SEQUENCE LISTING

<110> Mitchell, Lloyd G.
Garcia-Blanco, Mariano A.
Puttaraju, Madaiah
Mansfield, Gary S.

<120> METHODS AND COMPOSITIONS FOR USE IN
SPLICEROSOME MEDIATED RNA TRANS-SPLICING

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<141> 2001-01-08

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<210> 33
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<210> 34
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<223> Oligonucleotide primer complimentary to the beta HCG6 gene (accession #X00266)

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<220>

<223> Oligonucleotide primer complimentary to the Escherichia coli lacZ gene

<400> 37
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<211> 21

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<223> Oligonucleotide primer complimentary to the
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Escherichia coli lacZ gene

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<210> 42
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<212> DNA
<213> Homo sapiens

<400> 42
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<210> 43
<211> 51
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<210> 45
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<212> DNA
<213> Homo sapien

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aactagaagg cacagtcgag g 21

<210> 53
<211> 24
<212> DNA
<213> Artificial Sequence

<220>
<223> trans-spliced product containing Human chorionic
gonadotropin gene 6 sequences and Corynebacterium
diphtheriae diphtheria toxin A sequence

<400> 53
gagatgttcc agggcgtgat gatg 24

<210> 54
<211> 127
<212> RNA
<213> Artificial Sequence

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<223> PTM intramolecular base paired stem

<221> misc_feature
<222> (57)...(70)
<223> Loop comprising a combination of 14 nucleotides
according to specification

<400> 54

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nnnnnnnnnn aucguuaacu aauaaacuac uaacugggug aacuucuguu uuuuucucga 120
gcugcag 127

<210> 55
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<212> RNA
<213> Artificial Sequence

<220>
<223> PTM intramolecular base paired stem

<221> misc_feature
<222> (57)...(70)
<223> Loop comprising a combination of 14 nucleotides
according to specification

<400> 55
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nnnnnnnnnn aucguuaacu aauaaacuac uaacugggug aacuucugua uuauucucga 120
gcugcag 127

<210> 56
<211> 127
<212> RNA
<213> Artificial Sequence

<220>
<223> PTM intramolecular base paired stem

<221> misc_feature
<222> (57)...(70)
<223> Loop comprising a combination of 14 nucleotides
according to specification

<400> 56
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nnnnnnnnnn aucguuaacu aauaaacuac uaacugggug aaguucuguc cuugucucga 120
gcugcag 127

<210> 57
<211> 132
<212> DNA
<213> Artificial Sequence

<220>
<223> trans-spliced product containing Human chorionic
gonadotropin gene 6 sequences and Corynebacterium
diphtheriae diphtheria toxin A sequences

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tccattcaaa aa 132

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<213> Artificial Sequence

<220>
<223> Artificial Sequence derived from Escherichia coli
lacZ gene

<400> 58
gaattcggta ccatgggg 18

<210> 59
<211> 33
<212> DNA
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<220>
<223> Artificial Sequence derived from Escherichia coli
lacZ gene

<400> 59
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<210> 60
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<212> DNA
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<220>
<223> Artificial Sequence derived from Escherichia coli
lacZ gene

<400> 60
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<210> 61
<211> 25
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<220>
<223> trans-spliced product containing Escherichia coli
lacZ gene sequences and Human chorionic
gonadotropin gene 6 exon 2 sequences

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<211> 286
<212> DNA
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<220>
<223> trans-spliced product containing Escherichia coli
lacZ gene sequences

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acgggcaacc cgtggtcggc ttacggcggt gattttggcg atacgccaa cgatcgccag 240
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<213> Artificial Sequence

<220>
<223> trans-spliced product containing Escherichia coli
lacZ gene sequences

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gtaacagtct tggcggttc gctaaatact ggcaggcggt tcgtcagttat ccccggttac 120
aggggctgct gctgttgctg ctgctgagca tggcgggac atgggcattcc aaggagccac 180
ttcggccacg gtggcc 196

<210> 64
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<212> DNA
<213> Artificial Sequence

<220>
<223> trans-spliced product comprising cystic fibrosis
transmembrane regulator-derived sequences and His
tag sequence

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tgatgattat gggagaactg gagccttcag agggtaaaat taagcacagt ggaagaattt 180
cattctgttc tcagtttcc tggattatgc ctggcaccat taaagaaaat atcatcttg 240

gcggccgcca ctgtgctgga tatctgcaga attccaccac actggactag tggatccgag 300
ctcgttacca aggttaagtt taaaccgctg atcagcctcg actgtgcctt ctagttgccca 360
gccatctgtt gtttgccttccccgtgcc ttccttgacc ctggaagggtg ccactcccac 420

<210> 65
<211> 20
<212> DNA
<213> Artificial Sequence

<220>
<223> Splice junction sequence

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<210> 66
<211> 7
<212> PRT
<213> Artificial Sequence

<220>
<223> C terminal residues from glutathione -S- transferase

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<210> 67
<211> 15
<212> DNA
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<400> 67
ggagttgatc ccgtc 15

<210> 68
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<220>
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<223> Binding domain of PTM

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<220>
<223> Spacer sequence of PTM

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<210> 71
<211> 47
<212> DNA
<213> Artificial Sequence

<220>
<223> Branch point, pyrimidine tract and acceptor splice site of PTM

<400> 71
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<212> DNA
<213> Artificial Sequence

<220>
<223> Donor site and spacer sequence of PTM

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gatccaccgg 70

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<220>
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<210> 75
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<210> 76
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<210> 78
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ctagggttac cgaagtaaaa ccatacttat tag 33

<210> 79
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<220>
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<400> 79
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<210> 80
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<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide primer

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<210> 81
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<213> Artificial Sequence

<220>

<223> Binding domain of PTM molecule

<400> 81

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<210> 82

<211> 22

<212> DNA

<213> Artificial Sequence

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<223> Oligonucleotide primer

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<210> 83

<211> 21

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer

<400> 83

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<210> 84

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer

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<212> DNA

<213> Artificial Sequence

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<221> misc_feature

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<223> spacer sequence, see SEQ ID NO 70

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<211> 71
<212> DNA
<213> Artificial Sequence

<220>
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tatgtgaaa a 71

<210> 87
<211> 66
<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide

<400> 87
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acgccc 66

<210> 88
<211> 192
<212> DNA
<213> Artificial Sequence

<220>
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tccggccgca tcagctttg cagccaattc agttggatca tgcccggtac catcaaggag 120
aacataatct tcggcgtag ttacgacgag taccgctatc gctcggtgat taaggcctgt 180
cagttggagg ag 192

<210> 89
<211> 25
<212> DNA
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<220>
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<400> 89

gagcaggcaa gacgagcttg ctcat 25

<210> 90
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<212> DNA
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<220>
<223> Oligonucleotide

<400> 90
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<210> 91
<211> 30
<212> DNA
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<220>
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<400> 91
gtcagttgga ggaggacatc tccaaagtgg 30

<210> 92
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<220>
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tccggccgca tcagctttg cagccaattc agttggatca tgcccggtag catcaaggag 120
aacataatct tcggcgtag ttacgacgag taccgctatc gctcggtat taaggcctgt 180
cagttggagg ag 192

<210> 93
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<212> DNA
<213> Artificial Sequence

<220>
<223> PTM sequences

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aaatatcatt ggtgtttctt atgatga 27

<210> 94

<211> 30
<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide

<400> 94
ccaactagaa gaggacatct ccaagttgc 30

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<212> DNA
<213> Artificial Sequence

<220>
<223> Oligonucleotide

<400> 95
atgatcatgg gcgagttaga accaagttag 30

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<220>
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<400> 97
ccaactagaa gaggacatct ccaagtt 27

<210> 98
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<212> DNA
<213> Artificial Sequence

<220>
<223> 5' splice site

<400> 98
cgtttacagg taagtggatc c 21

<210> 99
<211> 27
<212> DNA
<213> Artificial Sequence

<220>
<223> 3' splice site

<400> 99
ctgcagggcg gcttcgtcta ataatgg 27

<210> 100
<211> 47
<212> DNA
<213> Artificial Sequence

<220>
<223> Sequence from trans-splicing domain

<400> 100
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<210> 101
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<212> DNA
<213> Artificial Sequence

<220>
<223> CFTR PTM

<400> 101
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ccttcgttg attctgctga caatctatct gaaaaattgg aaagagaatg ggatagagag 180
ctggcttcaa agaaaaatcc taaactcatt aatgcccttc ggcgatgtt tttctggaga 240
tttatgttct atggaatctt tttatattta ggggaagtca ccaaagcagt acagcctctc 300
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<210> 102
<211> 323
<212> DNA
<213> Artificial Sequence

<220>
<223> trans-splicing domain of CFTR PTM

<400> 102
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ctgtatctat attcatcatt ggaaacacca atgatatttt cttaatggt gcctggcata 180
atcctggaaa actgataaca caatgaaatt cttccactgt gcttaatttt accctctgaa 240
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ttaactcatt atcaaatcac gct 323

<210> 103
<211> 165
<212> DNA
<213> Artificial Sequence

<220>
<223> PTM binding domain

<400> 103
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cctaagcaga agtgtatatt cttatttgta aagattctat taactcattt gattcaaaat 120
atttaaaata cttccctgttt cacctactct gctatgcacc cgccgg 165

<210> 104
<211> 225
<212> DNA
<213> Artificial Sequence

<220>
<223> trans-splicing domain of CFTR PTM

<400> 104
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gcagaagtgt atatttttat ttgtaaagat tctattaact catttggattc aaaatattta 120
aaatacttcc tggcaccc actctgctat gcacccgcgg aacattatta taacgttgct 180
cgaatactaa ctggtaccc ttctttttt tttgatatcc tgtag 225

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<212> DNA
<213> Artificial Sequence

<220>
<223> CFTR PTM sequence

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aatatcatct ttgggtttc ctatgatgaa tatagataca gaagcgtcat caaagcatgc 180
caactagaag aggacatctc caagtttgcg gagaaagaca atatagttt tggagaaggt 240
ggaatcacac tgagtggagg tcaacgagca agaatttctt tagcaagagc agtatacaaa 300
gatgctgatt tgtattttt agactctcct tttggatacc tagatgttt aacagaaaaaa 360
gaaatatttg aaagctgtgt ctgtaaactg atggctaaca aaacttagat tttggtcact 420
tctaaaatgg aacattaaa gaaagctgac aaaatattaa ttttgcata agtagcagc 480
tatttttatg ggacatttc agaactccaa aatctacagc cagacttttgc ctcaaaactc 540
atggatgtg attcttcga ccaatttagt gcagaaagaa gaaattcaat cctaactgag 600
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caatctttt aacagactgg agagttggg gaaaaaaagga agaattctat tctcaatcca 720
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